

CECH et al.  
Application No.: 08/974,584  
Page 2

PATENT

a) Trp-R1-X7-R1-R1-R2-X-Phe-Phe-Tyr-X-Thr-Glu-X8-9-R3-R3-Arg-R4-X2-Trp

b) X3-Arg-X2-Pro-Lys-X3

c) X-Arg-X-Ile-X

d) X4-Phe-X3-Asp-X4-Tyr-Asp-X2

e) Tyr-X4-Gly-X2-Gln-Gly-X3-Ser-X8

f) X6-Asp-Asp-X-Leu-X3

wherein R1 is Leu or Ile; R2 is Gln or Arg; R3 is Phe or Tyr; <sup>R4 is Lys</sup>  
or His, and Xn represents the number n of consecutive unspecified amino acids;  
and wherein the protein has telomerase catalytic activity when  
complexed with a telomerase RNA component.

120. The polynucleotide of claim 119, encoding a protein that  
comprises the structure Trp-Leu-X-Tyr-X2-h-h-X-h-h-X-p-Phe-Phe-Tyr-X-Thr-Glu-X-  
p-X3-p-X3-Tyr-X-Arg-Lys-X2-Trp; wherein h is a hydrophobic amino acid selected  
from Ala, Leu, Ile, Val, Pro, Phe, Trp, and Met; and p is a polar amino acid selected  
from Gly, Ser, Thr, Tyr, Cys, Asn and Gln.

121. The polynucleotide of claim 119, where structure i) further  
comprises Arg-Lys-X2-Trp-X2-Leu.

122. The polynucleotide of claim 119, where structure b) comprises h-  
Arg-h-X-Pro-Lys, wherein h is a hydrophobic amino acid selected from Ala, Leu, Ile,  
Val, Pro, Phe, Trp, and Met.

123. The polynucleotide of claim 119, where structure c) comprises  
Arg-X-Ile-Pro-Lys.